

re-run



IFWO

RAW SEQUENCE LISTING

DATE: 10/15/2004

PATENT APPLICATION: US/10/702,319A

TIME: 10:59:38

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Output Set: N:\CRF4\10152004\J702319A.raw

1 <110> APPLICANT: Perera, Ranjan
 2 Rice, Stephen
 3 Eagleton, Clare
 4 <120> TITLE OF INVENTION: Compositions and Methods for the
 5 Modification of Gene Expression
 6 <130> FILE REFERENCE: 11000.1036c5
 C--> 7 <140> CURRENT APPLICATION NUMBER: US/10/702,319A
 8 <141> CURRENT FILING DATE: 2003-11-06
 9 <150> PRIOR APPLICATION NUMBER: U.S. No. 10/291,447
 10 <151> PRIOR FILING DATE: 2002-11-08
 11 <150> PRIOR APPLICATION NUMBER: U.S. No. 60/425,087
 12 <151> PRIOR FILING DATE: 2002-11-08
 13 <150> PRIOR APPLICATION NUMBER: U.S. No. 10/137,036
 14 <151> PRIOR FILING DATE: 2002-04-30
 15 <150> PRIOR APPLICATION NUMBER: U.S. No. 09/724,624
 16 <151> PRIOR FILING DATE: 2000-11-28
 17 <150> PRIOR APPLICATION NUMBER: U.S. No. 09/598,401
 18 <151> PRIOR FILING DATE: 2000-06-20
 19 <150> PRIOR APPLICATION NUMBER: PCT/NZ00/00018
 20 <151> PRIOR FILING DATE: 2000-02-24
 21 <150> PRIOR APPLICATION NUMBER: U.S. No. 60/146,591
 22 <151> PRIOR FILING DATE: 1999-07-30
 23 <150> PRIOR APPLICATION NUMBER: U.S. Patent No. 09/276,599
 24 <151> PRIOR FILING DATE: 1999-03-25
 25 <160> NUMBER OF SEQ ID NOS: 129
 26 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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 31 <213> ORGANISM: Pinus radiata
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 34 <222> LOCATION: (1)...(2064)
 35 <220> FEATURE:
 36 <221> NAME/KEY: intron
 37 <222> LOCATION: (1196)...(2033)
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P.6

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49	ttatcaatgg	aaaaatccat	ctaccaaaact	tacttttcaag	aaaatccaag	gattatagag	300
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54	aaaattttga	agtttaacaa	aaataacttg	gatctacaaa	aatccgtatc	ggattttctc	600
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107 <222> LOCATION: (1196) ... (2033)
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171      gggagaaggg tctcatccaa cgctattaaa tactcgccct caccgcgtta cttctcatct      1140
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191      ctctaactaa taaaacaata atcaccaaaa atctatcacc aaaaatgaaa aaagattttg      180
192      aatactaggc catatgagct acacaaattt caaaagtatc ttacacttat tacgcaccgc      240
193      gatgtcccca ctttcgaaaa acccgtttca agcctttcac gaaagtccaa cggtcagaaa      300
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201 <212> TYPE: DNA
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212      atagaaacaa ccagcaaagt tactagcagg aaatccaact aggtatcatg aagactacca      180
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243 <212> TYPE: DNA
244 <213> ORGANISM: Eucalyptus grandis
245 <220> FEATURE:
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:21; N Pos. 482

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L:444 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:480

L:2729 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (121) SEQUENCE:

L:2733 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (122) SEQUENCE:

L:2737 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (123) SEQUENCE:

L:2741 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (124) SEQUENCE: